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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/761,782

DATE: 08/07/2001
TIME: 13:21:02

Input Set : A:\202048US0.txt
Output Set: N:\CRF3\08072001\I761782.raw

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 3 <110> APPLICANT: LIVSHITS, VITALIY
 4         DOROSHENKO, VERA
 5         GORSHKOVA, NATALIYA
 6         BELARYEVA, ALLA
 7         IVANOVSKAYA, LIRINA
 8         KHOURGES, EVGENI
 9         AKHVERDIAN, VALERY
10         GUSYATINER, MIKHAIL
11         KOZLOV, YURY
13 <120> TITLE OF INVENTION: MUTANT ILVH GENE AND METHOD FOR PRODUCING L-VALINE
15 <130> FILE REFERENCE: 202048US0
17 <140> CURRENT APPLICATION NUMBER: 09/761,782
18 <141> CURRENT FILING DATE: 2001-01-18
20 <150> PRIOR APPLICATION NUMBER: RU 2000101678
21 <151> PRIOR FILING DATE: 2000-01-26
23 <160> NUMBER OF SEQ ID NOS: 8
25 <170> SOFTWARE: PatentIn version 3.1
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 492
29 <212> TYPE: DNA
30 <213> ORGANISM: Escherichia coli
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (1)..(489)
35 <223> OTHER INFORMATION:
38 <400> SEQUENCE: 1
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40 Met Arg Arg Ile Leu Ser Val Leu Leu Glu Asn Glu Ser Gly Ala Leu
41 1               5                   10                  15
43 tcc cgc gtg att ggc ctt ttt tcc cag cgt ggc tac aac att gaa agc      96
44 Ser Arg Val Ile Gly Leu Phe Ser Gln Arg Gly Tyr Asn Ile Glu Ser
45             20                  25                  30
47 ctg acc gtt gcg cca acc gac gat ccg aca tta tcg cgt atg acc atc     144
48 Leu Thr Val Ala Pro Thr Asp Asp Pro Thr Leu Ser Arg Met Thr Ile
49         35                  40                  45
51 cag acc gtg ggc gat gaa aaa gta ctt gag cag atc gaa aag caa tta     192
52 Gln Thr Val Gly Asp Glu Lys Val Leu Glu Gln Ile Glu Lys Gln Leu
53     50                  55                  60
55 cac aaa ctg gtc gat gtc ttg cgc gtg agt gag ttg ggg cag ggc gcg     240
56 His Lys Leu Val Asp Val Leu Arg Val Ser Glu Leu Gly Gln Gly Ala
57 65                  70                  75                  80
59 cat gtt gag cgg gaa atc atg ctg gtg aaa att cag gcc agc ggt tac     288
60 His Val Glu Arg Glu Ile Met Leu Val Lys Ile Gln Ala Ser Gly Tyr
61                 85                  90                  95
63 ggg cgt gac gaa gtg aaa cgt aat acg gaa ata ttc cgt ggg caa att     336
64 Gly Arg Asp Glu Val Lys Arg Asn Thr Glu Ile Phe Arg Gly Gln Ile
65                 100                 105                 110
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67 atc gat gtc aca ccc tcg ctt tat acc gtt caa tta gca ggc acc agc      384
68 Ile Asp Val Thr Pro Ser Leu Tyr Thr Val Gln Leu Ala Gly Thr Ser
69         115                 120                 125
71 ggt aag ctt agt gca ttt tta gca tcg att cgc gat gtg gcg aaa att      432
72 Gly Lys Leu Ser Ala Phe Leu Ala Ser Ile Arg Asp Val Ala Lys Ile
73         130                 135                 140
75 gtg gag gtt gct cgc tct ggt gtg gtc gga ctt tcg cgc ggc gat aaa      480
76 Val Glu Val Ala Arg Ser Gly Val Val Gly Leu Ser Arg Gly Asp Lys
77 145                 150                 155                 160
79 ata atg cgt tga                                                      492
80 Ile Met Arg
84 <210> SEQ ID NO: 2
85 <211> LENGTH: 163
86 <212> TYPE: PRT
87 <213> ORGANISM: Escherichia coli
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91 Met Arg Arg Ile Leu Ser Val Leu Leu Glu Asn Glu Ser Gly Ala Leu
92 1               5                   10                  15
95 Ser Arg Val Ile Gly Leu Phe Ser Gln Arg Gly Tyr Asn Ile Glu Ser
96         20                  25                  30
99 Leu Thr Val Ala Pro Thr Asp Asp Pro Thr Leu Ser Arg Met Thr Ile
100         35                  40                  45
103 Gln Thr Val Gly Asp Glu Lys Val Leu Glu Gln Ile Glu Lys Gln Leu
104        50                  55                  60
107 His Lys Leu Val Asp Val Leu Arg Val Ser Glu Leu Gly Gln Gly Ala
108 65                  70                  75                  80
111 His Val Glu Arg Glu Ile Met Leu Val Lys Ile Gln Ala Ser Gly Tyr
112                  85                  90                  95
115 Gly Arg Asp Glu Val Lys Arg Asn Thr Glu Ile Phe Arg Gly Gln Ile
116              100                 105                 110
119 Ile Asp Val Thr Pro Ser Leu Tyr Thr Val Gln Leu Ala Gly Thr Ser
120         115                 120                 125
123 Gly Lys Leu Ser Ala Phe Leu Ala Ser Ile Arg Asp Val Ala Lys Ile
124        130                 135                 140
127 Val Glu Val Ala Arg Ser Gly Val Val Gly Leu Ser Arg Gly Asp Lys
128 145                 150                 155                 160
131 Ile Met Arg
135 <210> SEQ ID NO: 3
136 <211> LENGTH: 18
137 <212> TYPE: DNA
138 <213> ORGANISM: Artificial Sequence
140 <220> FEATURE:
141 <223> OTHER INFORMATION: Synthetic DNA
143 <400> SEQUENCE: 3
144 gacatgaatg tctggttt                                                  18
147 <210> SEQ ID NO: 4
148 <211> LENGTH: 19
149 <212> TYPE: DNA
150 <213> ORGANISM: Artificial Sequence
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153 <223> OTHER INFORMATION: Synthetic DNA
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159 <210> SEQ ID NO: 5
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161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial Sequence
164 <220> FEATURE:
165 <223> OTHER INFORMATION: Synthetic DNA
167 <400> SEQUENCE: 5
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172 <211> LENGTH: 24
173 <212> TYPE: DNA
174 <213> ORGANISM: Artificial Sequence
176 <220> FEATURE:
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184 <211> LENGTH: 25
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
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196 <211> LENGTH: 27
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198 <213> ORGANISM: Artificial Sequence
200 <220> FEATURE:
201 <223> OTHER INFORMATION: Synthetic DNA
203 <400> SEQUENCE: 8
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VERIFICATION SUMMARY   DATE: 08/07/2001
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